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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

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Revised 10/08/03



DATE: 12/04/2003

IFWO

PATENT APPLICATION: US/10/716,480 TIME: 10:54:22 Input Set : N:\EBONY'S\10716480.raw.txt Output Set: N:\CRF4\12042003\J716480.raw 3 <110> APPLICANT: Ajinomoto Co. Inc. 5 <120> TITLE OF INVENTION: Method for Producing L-Amino Acid Using Methylotroph 7 <130> FILE REFERENCE: OP1627/US-102 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/716,480 C--> 10 <141> CURRENT FILING DATE: 2003-11-20 12 <150> PRIOR APPLICATION NUMBER: JP 2002-336315 Does Not Comply 13 <151> PRIOR FILING DATE: 2002-11-20 Corrected Diskette Needed 15 <160> NUMBER OF SEQ ID NOS: 14 17 <170> SOFTWARE: PatentIn Ver. 2.0 ERRORED SEQUENCES 252 <210> SEQ ID NO: 739 Shand 253 <211> LENGTH: (33) 254 <212> TYPE: DNA 255 <213> ORGANISM: Artificial Sequence 257 <220> FEATURE: 258 <223> OTHER INFORMATION: Description of Artificial Sequence: primer 260 <400> SEQUENCE: 7 E--> 261 agggaattcc ccgttctgga taatgttttt tgcgccgac 286 <210> SEQ ID NO: 10. 287 <211> LENGTH: (33)-50 Sh(W) 288 <212> TYPE: DNA 289 <213> ORGANISM: Artificial Sequence 291 <220> FEATURE: 292 <223> OTHER INFORMATION: Description of Artificial Sequence: primer 294 <400> SEQUENCE: 10 E--> 295 gggggageta gaagagetee aaaacccgcg aaaactaacc catcaacate 50 330 <210> SEQ ID NO: 14 331 <211> LENGTH: 32 332 <212> TYPE: DNA 333 <213> ORGANISM: Artificial Sequence 335 <220> FEATURE: 336 <223> OTHER INFORMATION: Description of Artificial Sequence: primer 338 <400> SEQUENCE: 14 32 339 gccgtgttgc taggatggtt gttcttggat ca E--> 343(2)

RAW SEQUENCE LISTING

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/716,480

DATE: 12/04/2003 TIME: 10:54:23

Input Set : N:\EBONY'S\10716480.raw.txt
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L:9 M:270 C: Current Application Number differs, Replaced Application Number L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:261 M:252 E: No. of Seq. differs, <211> LENGTH:Input:33 Found:39 SEQ:7 L:295 M:252 E: No. of Seq. differs, <211> LENGTH:Input:33 Found:50 SEQ:10 L:343 M:254 E: No. of Bases conflict, this line has no nucleotides.